

-120

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1

FEVGGAAPFNRTHRSKRSSSHPIFHRGEFSVCDSVSVWVGDKTTATDIKGKE

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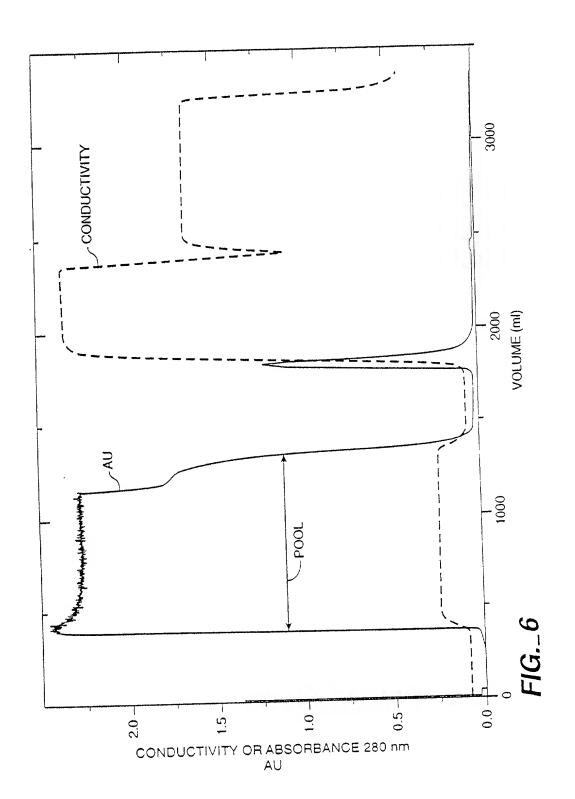
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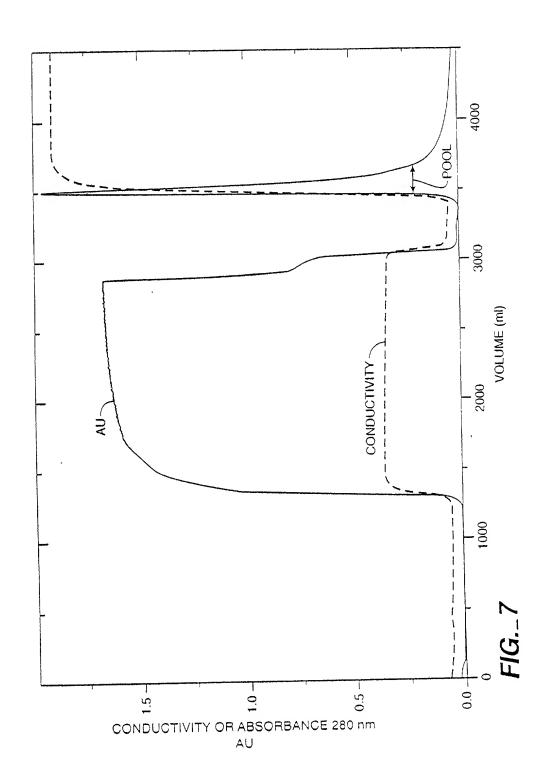
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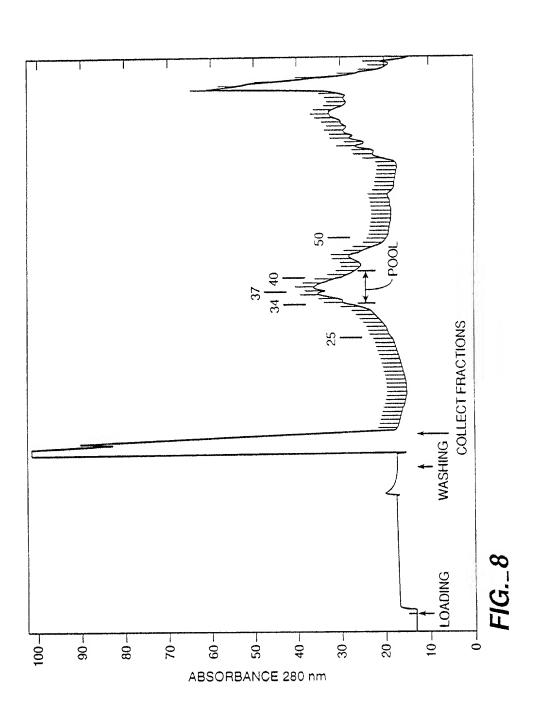
FIG._4

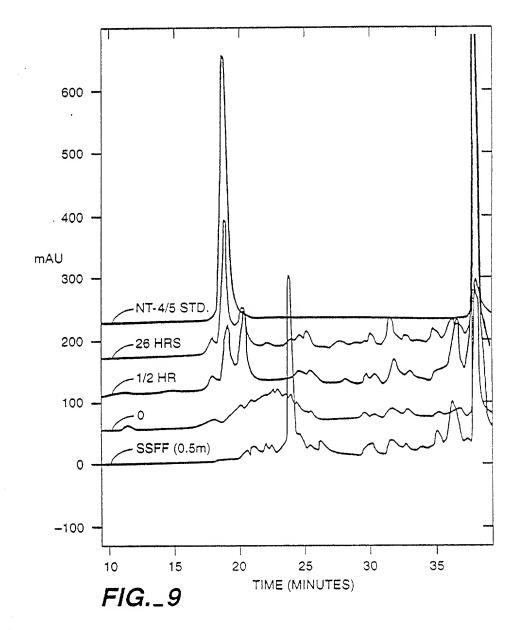
	_	
rhNGF	1	SSSHPIFHRGEFSVCDSVSVWVGDKTTATDIKGKEVMVLGEVN-IN
muNGF	. 1	SSTHPVFHMGEFSVCDSVSVWVGDKTTATDIKGKEVTVLAEVN-IN
BDNF	1	HSDPA-RRGELSVCDSISEWVTAADKKTAVDMSGGTVTVLEKVP-VS
NT-3	1	YAEHKSHRGEYSVCDSESLWVTDKSSAIDIRGHQVTVLGEIK-TG
NT-4/5	1	GVSETAPASRRGELAVCDAVSGWVTDRRTAVDLRGREVEVLGEVPAAG
rhngf	46	NSVFKQYFFETKCRDPNPVDSGCRGIDSKHWNSYCTTTHTFVK
muNGF	46	NSVFRQYFFETK CRASNPVESGC RGIDSKHWNSYCTTTHTFVK
BDNF	46	KGQLKQYFYETKCNPMGYTKEGCRGIDKRHWNSQCRTTQSYVR
NT-3	45	NSPVKQYFYETRCKEARPVKNGCRGIDDKHWNSQCKTSQTYVR
NT-4/5	49	GSPLRQYFFETRCKADNAEEGGPGAGGGGCRGVDRRHWVSECKAKQSYVR
rhNGF	89	ALTMD-GKQAAWRFIRIDTACVCVLSRKAVRRA
muNGF	89	ALTTD-EKQAAWRFIRIDTACVCVLSRKATRRG
BDNF	89	ALTMDSKKRIGWRFIRIDTSCVCTLTIKRGR
NT-3	88	ALTSENNKLVGWRWIRIDTSCVSALSRKIGRT-
NT-4/5	99	ALTAHAQGRVGWRWIRIDTACVCTLLSRTGRA-

FIG._5









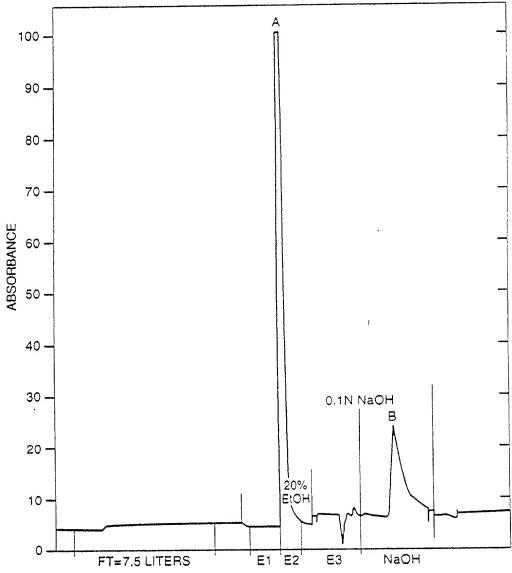
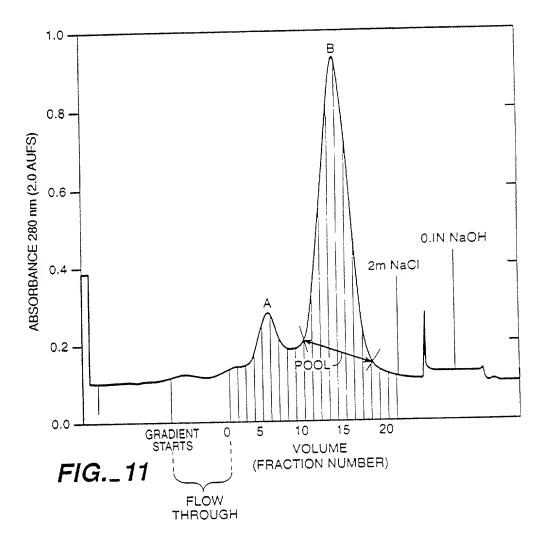
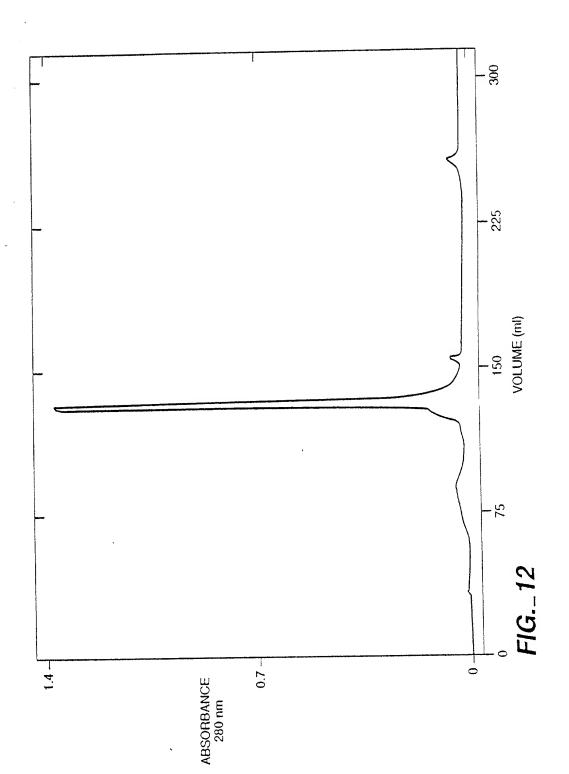


FIG._10





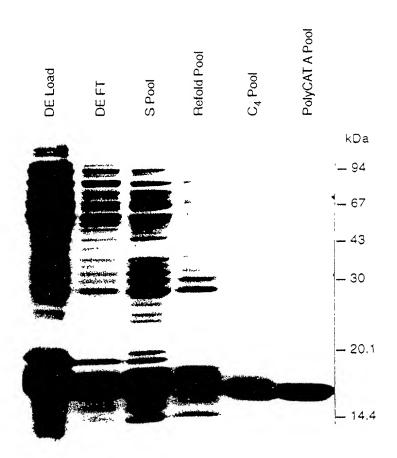


FIG._13

